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SERIAL NUMBER	FILING DATE	FIRST NAMED APPLICANT	ATTORNEY DOCKET NO

EXAMINER	
ART UNIT	PAPER NUMBER

DATE MAILED:

Please find below a communication from the EXAMINER in charge of this application

Commissioner of Patents

This application contains sequence disclosures that are encompassed by the definitions for nucleotide and/or amino acid sequences set forth in 37 CFR 1.821(a)(1) and (a)(2). However, this application fails to comply with the requirements of 37 CFR 1.821 through 1.825 for the reason(s) set forth on the attached Notice To Comply With Requirements For Patent Applications Containing Nucleotide Sequence And/Or Amino Acid Sequence Disclosures and marked-up Raw Sequence Listing.

APPLICANT MUST COMPLY WITH THE SEQUENCE RULES, 37 CFR 1.821 - 1.825 IN THE TIME GIVEN FOR RESPONSE TO THIS OFFICE ACTION. Failure to comply with these requirements will result in ABANDONMENT of the application under 37 CFR 1.821(g). Extensions of time may be obtained by filing a petition accompanied by the extension fee under the provisions of 37 CFR 1.136. In no case may an applicant extend the period for response beyond the six month statutory period. Applicant is requested to return a copy of the attached Notice to Comply with the response.

Any inquiry concerning this communication or earlier communications from the Examiner should be directed to Holly Schnizer, Ph.D. whose telephone number is (703) 305-3722. The Examiner can normally be reached Monday through Thursday from 7:30 AM to 4:30 PM.

If attempts to reach the Examiner by telephone are unsuccessful, the Examiner's supervisor, Christopher Low, can be reached at (703) 306-4119. The OFFICIAL fax phone number for Technology Center 1600 is (703) 308-4242.

Any inquiry of a general nature or relating to the status of this application or proceeding should be directed to the Technology Center 1600 receptionist whose telephone number is (703) 308-0196.

**NOTICE TO COMPLY WITH REQUIREMENTS FOR PATENT APPLICATIONS CONTAINING
NUCLEOTIDE SEQUENCE AND/OR AMINO ACID SEQUENCE DISCLOSURES**

The nucleotide and/or amino acid sequence disclosure contained in this application does not comply with the requirements for such a disclosure as set forth in 37 C.F.R. 1.821 - 1.825 for the following reason(s):

- 1. This application clearly fails to comply with the requirements of 37 C.F.R. 1.821-1.825. Applicant's attention is directed to these regulations, published at 1114 OG 29, May 15, 1990 and at 55 FR 18230, May 1, 1990.
- 2. This application does not contain, as a separate part of the disclosure on paper copy, a "Sequence Listing" as required by 37 C.F.R. 1.821(c).
- 3. A copy of the "Sequence Listing" in computer readable form has not been submitted as required by 37 C.F.R. 1.821(e).
- 4. A copy of the "Sequence Listing" in computer readable form has been submitted. However, the content of the computer readable form does not comply with the requirements of 37 C.F.R. 1.822 and/or 1.823, as indicated on the attached copy of the marked -up "Raw Sequence Listing".
- 5. The computer readable form that has been filed with this application has been found to be damaged and/or unreadable as indicated on the attached CRF Diskette Problem Report. A Substitute computer readable form must be submitted as required by 37 C.F.R. 1.825(d).
- 6. The paper copy of the "Sequence Listing" is not the same as the computer readable form of the "Sequence Listing" as required by 37 C.F.R. 1.821(e).
- 7. Other: _____

Applicant Must Provide:

- An initial or substitute computer readable form (CRF) copy of the "Sequence Listing".
- An initial or substitute paper copy of the "Sequence Listing", as well as an amendment directing its entry into the specification.
- A statement that the content of the paper and computer readable copies are the same and, where applicable, include no new matter, as required by 37 C.F.R. 1.821(e) or 1.821(f) or 1.821(g) or 1.825(b) or 1.825(d).

For questions regarding compliance to these requirements, please contact:

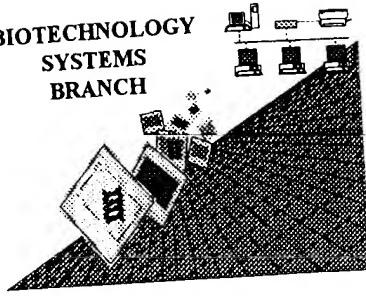
For Rules Interpretation, call (703) 308-4216
For CRF Submission Help, call (703) 308-4212
For PatentIn software help, call (703) 308-6856

PLEASE RETURN A COPY OF THIS NOTICE WITH YOUR RESPONSE

Schnizer

RAW SEQUENCE LISTING ERROR REPORT

BIOTECHNOLOGY
SYSTEMS
BRANCH



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following CRF diskette:

Application Serial Number:

09/185,904

Art Unit / Team No.:

1653

Date Processed by STIC:

12/10/99

BEST AVAILABLE COPY

**THE ATTACHED PRINTOUT EXPLAINS THE ERRORS DETECTED.
PLEASE BE SURE TO FORWARD THIS INFORMATION TO THE APPLICANTS
BY EITHER:**

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANTS ALONG WITH A NOTICE TO COMPLY or,
- 2) CALLING APPLICANTS AND FAXING THEM A COPY OF THE PRINTOUT WITH A NOTICE TO COMPLY

THIS WILL INSURE THAT THE NEXT SUBMISSION RECEIVED FROM THEM WILL BE ERROR FREE.

IF YOU HAVE ANY FURTHER QUESTIONS, PLEASE CALL:

MARK SPENCER 703-308-4212

Raw Sequence Listing Error Summary

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER: 09/185,904

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleics The number/text at the end of each line "wrapped" down to the next line.
This may occur if your file was retrieved in a word processor after creating it.
Please adjust your right margin to .3, as this will prevent "wrapping".
- 2 Wrapped Aminos The amino acid number/text at the end of each line "wrapped" down to the next line.
This may occur if your file was retrieved in a word processor after creating it.
Please adjust your right margin to .3, as this will prevent "wrapping".
- 3 Incorrect Line Length The rules require that a line not exceed 72 characters in length. This includes spaces.
- 4 Misaligned Amino Acid Numbering The numbering under each 5th amino acid is misaligned. This may be caused by the use of tabs between the numbering. It is recommended to delete any tabs and use spacing between the numbers.
- 5 Non-ASCII This file was not saved in ASCII (DOS) text, as required by the Sequence Rules.
Please ensure your subsequent submission is saved in ASCII text so that it can be processed.
- 6 Variable Length Sequence(s) _____ contain n's or Xaa's which represented more than one residue.
As per the rules, each n or Xaa can only represent a single residue.
Please present the maximum number of each residue having variable length and indicate in the (ix) feature section that some may be missing.
- 7 PatentIn ver. 2.0 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequence(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence.
- 8 Skipped Sequences (OLD RULES) Sequence(s) _____ missing. If intentional, please use the following format for each skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X:
(i) SEQUENCE CHARACTERISTICS:(Do not insert any headings under "SEQUENCE CHARACTERISTICS")
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X:
This sequence is intentionally skipped

Please also adjust the "(iii) NUMBER OF SEQUENCES:" response to include the skipped sequence(s).
- 9 Skipped Sequences (NEW RULES) Sequence(s) _____ missing. If intentional, please use the following format for each skipped sequence.
<210> sequence id number
<400> sequence id number
000
- 10 Use of n's or Xaa's (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
Use of <220> to <223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 11 Use of <213>Organism (NEW RULES) Sequence(s) _____ are missing this mandatory field or its response.
5-9, 11, 13, 15-17, 19-23, 25, 27-29 (maybe more)
- 12 Use of <220>Feature (NEW RULES) Sequence(s) _____ are missing the <220>Feature and associated headings.
Use of <220> to <223> is MANDATORY if <213>ORGANISM is "Artificial" or "Unknown"
Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 6/01/98, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of new Rules)
- 13 PatentIn ver. 2.0 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other means to copy file to floppy disk.

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RAW SEQUENCE LISTING
PATENT APPLICATION US/09/185,904

DATE: 12/10/1999
 TIME: 15:36:57

Input Set: I185904.RAW

**This Raw Listing contains the General Information
Section and up to first 5 pages.**

1 <110> APPLICANT: Anderson, Christen M.
 2 Davis, Robert E.
 3 Clevenger, William
 4 Wiley, Sandra Eileen
 5 Willer, Scott W.
 6 Szabo, Tomas R.
 7 Ghosh, Soumitra S.

Does Not Comply
Corrected Diskette Needed
pp 2-5

8 <120> TITLE OF INVENTION: PRODUCTION OF ADENINE NUCLEOTIDE
 9 TRANSLOCATOR (ANT), NOVEL ANT LIGANDS AND SCREENING ASSAYS
 10 THEREFOR

11 <130> FILE REFERENCE: 660088.420

12 <140> CURRENT APPLICATION NUMBER: US/09/185,904

13 <141> CURRENT FILING DATE: 1998-11-03

14 <160> NUMBER OF SEQ ID NOS: 33

15 <170> SOFTWARE: FastSEQ for Windows Version 3.0

16 <210> SEQ ID NO 1

17 <211> LENGTH: 894

18 <212> TYPE: DNA

19 <213> ORGANISM: Homo sapien

20 <400> SEQUENCE: 1

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30 caaggcatca ttatctatac agctgcctac ttggaggtct atgataactgc caagggatg	600
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43 gccagcaagc agatcactgc agataagcaa tacaaaggca ttataactg cgtggccgt	180
44 atcccaagg agcaggaagt tcttccttc tggcgccgtt acctggccaa tgtcatcaga	240

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RAW SEQUENCE LISTING
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47 ggtggcccgag gggccacatc cctgtgttt gtgtaccctc ttgattttgc cctgtaccctg 420
48 ctagcagctg atgtgggtaa agctggagct gaaaqqqaat tccgaggcct cggtgactgc 480
49 ctgggttaaga tctacaaatc tgatggatt aagggcctgt accaaggctt taacgtgtct 540
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53 cagtcagggtt gcaaaggaaac tgacatcatg tacacaggca cgcttgactg ctggcggaaag 780
54 attgctcgtt atgaaggagg caaagcttt ttcaagggtt catggtccaa tgttctcaga 840
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63 gccagcaAGC agatcgccgc cgacaacgcg tacaaggggca tcgtgactg cattgtccgc 180
64 atcccaagg agcagggtgt gctgtccctc tggaggggca accttgcCAA cgtcattcgc 240
65 tacttcccc CTCAGGCCCT caacttcgccc ttcaaggata agtacaagca gatttccctg 300
66 gggggcgtgg acaagcacac gcagttctgg aggtactttg cggcaacct ggcctccggc 360
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74 atcttcagag atgagggggg caaggccttc ttcaagggtt cgtggtccaa cgtcctgcgg 840
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92 <212> TYPE: DNA
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RAW SEQUENCE LISTING
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171 <213> ORGANISM: Artificial Sequence
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RAW SEQUENCE LISTING
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VERIFICATION SUMMARY
PATENT APPLICATION US/09/185,904

DATE: 12/10/1999
TIME: 15:36:57

Input Set: I185904.RAW

Line ? Error/Warning

Original Text